

AJ

SEQUENCE LISTING



<110> Burch, Ronald
Sackler, David

<120> Contraceptive Antibody Vaccines

<130> 6750-018-999

<140> 09/831,631
<141> 2001-05-10

<160> 70

<170> PatentIn version 3.0

<210> 1
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<220>
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<223> Description of Artificial Sequence: Primer for PCR

<400> 1
aacagctatg accatg 16

<210> 2
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<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Primer for PCR

<400> 2
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<210> 3
<211> 13
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<220>
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<223> Description of Artificial Sequence: CDR Derived peptide with Biotin label at the N-terminal residue

<400> 3
Thr Ala Lys Ala Ser Gln Ser Val Ser Asn Asp Val Ala
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<210> 4
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: CDR Derived peptide with Biotin

label at the N-terminal residue

<400> 4
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<210> 5
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: CDR Derived peptide with Biotin
label at the N-terminal residue

<400> 5
Phe Ala Gln Gln Asp Tyr Ser Ser Pro Leu Thr
1 5 10

<210> 6
<211> 7
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<220>
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<223> Description of Artificial Sequence: CDR Derived peptide with Biotin
label at the N-terminal residue

<400> 6
Phe Thr Asn Tyr Gly Met Asn
1 5

<210> 7
<211> 19
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<220>
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<223> Description of Artificial Sequence: CDR Derived peptide with Biotin
label at the N-terminal residue

<400> 7
Ala Gly Trp Ile Asn Thr Tyr Thr Gly Glu Pro Thr Tyr Ala Asp
1 5 10 15
Asp Phe Lys Gly

<210> 8
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: CDR Derived peptide with Biotin
label at the N-terminal residue

<400> 8
Ala Arg Ala Tyr Tyr Gly Lys Tyr Phe Asp Tyr
1 5 10

<210> 9
 <211> 221
 <212> DNA
 <213> Artificial Sequence

<220>
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 <223> Description of Artificial Sequence: Sperm cell specific epitope

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 agcacatcta caggcacaat attaaattgc tacacatgtg cttatatgaa tgatcaagga 180
 aaatgtcttc gtggagaggg aacctgcatc actcagaatt c 221

<210> 10
 <211> 69
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> misc_feature
 <223> Description of Artificial Sequence: Sperm cell specific epitope

<400> 10
 Gln Pro Ser Gly Glu His Gly Glu Gln Pro Ser Gly Glu Gln Ala Ser
 1 5 10 15
 Gly Glu Gln Pro Ser Gly Glu His Ala Ser Gly Glu Gln Ala Ser Gly
 20 25 30
 Ala Gln Ile Ser Ser Thr Ser Thr Gly Thr Ile Leu Asn Cys Tyr Thr
 35 40 45
 Cys Ala Tyr Met Asn Asp Gln Gly Lys Cys Leu Arg Gly Glu Gly Thr
 50 55 60
 Cys Ile Thr Gln Asn
 65

<210> 11
 <211> 75
 <212> DNA
 <213> Artificial Sequence

<220>
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 <223> Description of Artificial Sequence: Cloning primers for SP10

<400> 11
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 ggtgaacagc cttag 75

<210> 12
 <211> 75
 <212> DNA
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<220>
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 <223> Description of Artificial Sequence: Cloning primers for SP10

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<210> 13
 <211> 70
 <212> DNA
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 <223> Description of Artificial Sequence: Cloning primers for SP10

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 ctcagaattc 70

 <210> 14
 <211> 70
 <212> DNA
 <213> Artificial Sequence

 <220>
 <221> misc_feature
 <223> Description of Artificial Sequence: Cloning primers for SP10

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 ctcagaattc 70

 <210> 15
 <211> 79
 <212> DNA
 <213> Artificial Sequence

 <220>
 <221> misc_feature
 <223> Description of Artificial Sequence: Cloning primers for SP10

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 gcacatgtgt agcaattta 79

 <210> 16
 <211> 79
 <212> DNA
 <213> Artificial Sequence

 <220>
 <221> misc_feature
 <223> Description of Artificial Sequence: Cloning primers for SP10

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 gctgctgtgt agcaattta 79

 <210> 17
 <211> 75
 <212> DNA
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 <220>
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 <223> Description of Artificial Sequence: Cloning primers for SP10

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<210> 18
<211> 67
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description of Artificial Sequence: Cloning primers for SP10

<400> 18
gttctcccga ggctgtctca ccagaaggct gttcacccga gccatgttca cctgaaggct 60
ggaattc 67

<210> 19
<211> 210
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description of Artificial Sequence: Sperm cell specific epitope M
SA-6

<400> 19
gtcggcagcc tccgaagcag cccgctccag agcccgtgc tccgaccgct cgtccagagc 60
agcctctgct tgctgttctt cttgctgcga tacagctgcg gcgacggcag ctgcagccga 120
cgatactgcg acttgacggt gtgccggcga atgtacttgc tgctgcgatt cacggaccgc 180
ccgctcccgc agacgtgctg cgtcttgagc 210

<210> 20
<211> 70
<212> PRT
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description of Artificial Sequence: Sperm cell specific epitope M
SA-6

<400> 20
Gln Pro Ser Glu Ala Ser Ser Gly Glu Val Ser Gly Asp Glu Ala Gly
1 5 10 15
Glu Gln Val Ser Ser Glu Thr Asn Asp Lys Glu Asn Asp Ala Met Ser
20 25 30
Thr Pro Leu Pro Ser Thr Ser Ala Ala Ile Thr Leu Asn Cys His Thr
35 40 45
Cys Ala Tyr Met Asn Asp Asp Ala Lys Cys Leu Arg Gly Glu Gly Val
50 55 60
Cys Thr Thr Gln Asn Ser
65 70

<210> 21
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Oligomer from MSA 63

<400> 21
gtcggcagcc tccgaagcag cccgctccag agcccgtgc tccga 45

<210> 22
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Oligomer from MSA 63

<400> 22
agcccgtgc tccgaccgct cgtccagagc agcctctgct tgctg 45

<210> 23
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Oligomer from MSA 63

<400> 23
agcctctgct tgctgttccct cttgctgcga tacagctgcg gcgac 45

<210> 24
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Oligomer from MSA 63

<400> 24
tacagctgcg gcgacggcag ctgcagccga cgatactgcg acttg 45

<210> 25
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Oligomer from MSA 63

<400> 25
cgatactgcg acttgacggt gtgcacgcga atgtacttgc tgctg 45

<210> 26
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Oligomer from MSA 63

<400> 26
atgtacttgc tgctgcgatt cacggacgcg ccgctcccgc agacg 45

<210> 27
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Oligomer from MSA 63

<400> 27
cgattcacgg acgcgccgct cccgcagacg tgctgcgtct tgagc 45

<210> 28
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description of Artificial Sequence: Consensus sequence

<400> 28
Gln Pro Ser Glu Ala Ser Ser Gly Glu Val Ser Gly Asp Glu Ala Gly
1 5 10 15
Glu

<210> 29
<211> 384
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Consensus sequence

<400> 29
atggccttggg gttggacctt gctattcctg atggcagctg cccaaagtgc ccaagcagat 60
atgcaaataa cacaagtcc tagtagtttg agtgctagtg tgggagatca agtgacaatc 120
acatgtcggg ctagtcaaag tatcagtaac tgtttggcctt ggtatcaaca aaagcctgga 180
aaggctccta agttgttgat ctatgctgct agtagtttg agagtggagt gcctagtcgg 240
ttcagtggaa gtggaagtgg aacacggttc accttgacca tgagttagttt gcaacctgag 300
gatttcgcta cctattattg tcaacaatat aacagtttgc cttggacctt cggacaagga 360
accaaggtgg agatcaagga attc 384

<210> 30
<211> 417
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Consensus sequence

<400> 30
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gttcagctgg tgcagtgtgg cgctgaggtg aagaagcctg gcgcttctgt gaaggtgtct 120
tgcaaggctt ctggctacac attcacatct tacgctatat cttggaattg ggtgaggcag 180

gctcctggcc	agggcctgga	gtggatgggc	tggataaatg	gaaatggaga	tacaaattac	240
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atggagctgt	cttctctgag	gtctgaggat	actgctgttt	actactgcgc	tagggctcct	360
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<210> 31
 <211> 63
 <212> DNA
 <213> Artificial Sequence

<220>
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 <223> Description of Artificial Sequence: Consensus sequence

<400> 31	
gacattgtga	60
tgacacagtc	
tccatcctcc	
ctagctgtgt	
cagttggaga	
gaaggttact	63
atg	

<210> 32
 <211> 74
 <212> DNA
 <213> Artificial Sequence

<220>
 <221> misc_feature
 <223> Description of Artificial Sequence: Consensus sequence

<400> 32	
gcaagctcat	60
agtaaccttc	
tctccaactg	
acacacgata	
gggaggatgg	
agactgtgac	74
atcacaatgt	
ctgc	

<210> 33
 <211> 84
 <212> DNA
 <213> Artificial Sequence

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 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

<400> 33	
agctgcgtcg	60
gcagcctccg	
aagcagcccg	
ctccagagcc	
cgctgctggc	
atggtaccag	84
cagaaaccag	
ggcagctctc	
taaa	

<210> 34
 <211> 72
 <212> DNA
 <213> Artificial Sequence

<220>
 <221> misc_feature
 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

<400> 34	
ctgccctggg	60
ttctgctggg	
accatcggag	
cagcgggctc	
tgccggagcgg	
gctgcttcgg	72
acggctgccg	
ac	

<210> 35
 <211> 78

<212> DNA
 <213> Artificial Sequence

 <220>
 <221> misc_feature
 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

 <400> 35
 gacattgtga tgtcacagtc tccatcctcc ctagctgtgt cagttggaga gaagggttact 60
 gtgagcgcta agtccagt 78

 <210> 36
 <211> 75
 <212> DNA
 <213> Artificial Sequence

 <220>
 <221> misc_feature
 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

 <400> 36
 gagagccttt tatatagtag caatcaaaag atctacttgg cctggtacca gcagaaacca 60
 gggcagtctc ctaaa 75

 <210> 37
 <211> 67
 <212> DNA
 <213> Artificial Sequence

 <220>
 <221> misc_feature
 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

 <400> 37
 ctgctgattt actgggcac cactagggaa tctgggggtcc ctgatcgctt cacaggctgg 60
 atctggg 67

 <210> 38
 <211> 68
 <212> DNA
 <213> Artificial Sequence

 <220>
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 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

 <400> 38
 gcacagcaat attatagata tctcggacgt tcggtggagc caccaagctg caaatcaaac 60
 cggaattc 68

 <210> 39
 <211> 69
 <212> DNA
 <213> Artificial Sequence

 <220>
 <221> misc_feature

<223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

<400> 39
accgcctgtg aagcgatcag gcaccccgaga ttccctagtg gatgcccgagt aaatcagcag 60
ttaggaga 69

<210> 40
<211> 77
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

<400> 40
ctgccctggt ttctgctggt accaggccaa gtagatcttt tgagattgct actatataaa 60
aggctctgac tggactt 77

<210> 41
<211> 78
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

<400> 41
agcgctcata gtaaccttct ctccaactga cacagctagc gacgatcgag actgtgacat 60
cacaatgtct gcttgggc 78

<210> 42
<211> 78
<212> DNA
<213> Artificial Sequence

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<223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

<400> 42
gaattcccggt ttgatttcca gcttggtgcc tccaccgaac gtccgaggat atctataata 60
ttgctgtgcg taataaac 78

<210> 43
<211> 57
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

<400> 43
 agagatttga gtctgaccat cagcagtgtg aaggctgaag acgtggcagt ttattac 57

 <210> 44
 <211> 57
 <212> DNA
 <213> Artificial Sequence

 <220>
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 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

 <400> 44
 tgccagggtct tcagccttga cactgctgat ggtgagagtg aaatctgtcc cagatcc 57

 <210> 45
 <211> 66
 <212> DNA
 <213> Artificial Sequence

 <220>
 <221> misc_feature
 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

 <400> 45
 tcgtgccagt tcctcgtcga ctagctcttc gactagctcc tgctgctctt gtcggtcacg 60
 gaattc 66

 <210> 46
 <211> 75
 <212> DNA
 <213> Artificial Sequence

 <220>
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 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

 <400> 46
 gaattccgtg accgacaaga gcagcaggag ctagtcgaag agctggtcga cgaggaactg 60
 gcacgacggg ttcgt 75

 <210> 47
 <211> 80
 <212> DNA
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 <223> Description for Artificial Sequence: Constructs for LDH-C4

 <400> 47
 gaattcatgg cttgggtgtg gaccttgcta ttcctgatgg cagctgccca aagtgcccaa 60
 gcacagatcc agttgggtgca 80

 <210> 48
 <211> 79

<212> DNA
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<220>
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 <223> Description for Artificial Sequence: Constructs for LDH-C4

<400> 48
 gtctggacct gagctgaaga agcctggaga gacagtcaag atctccgcta aggcttctgg 60
 gtataccttc acaaactag 79

<210> 49
 <211> 80
 <212> DNA
 <213> Artificial Sequence

<220>
 <221> misc_feature
 <223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

<400> 49
 gaatgaactg ggtgaagcag gctccaggaa agggttttaa gtggatgggc tggataaaca 60
 cctacactgg agagccaaca 80

<210> 50
 <211> 80
 <212> DNA
 <213> Artificial Sequence

<220>
 <221> misc_feature
 <223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

<400> 50
 tatgctgatg acttcaaggg acggtttgcc ttctctttgg aaacctctgc cagcactgcc 60
 tatttgcaag atcaacacct 80

<210> 51
 <211> 70
 <212> DNA
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 <223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

<400> 51
 caaaaatgag gacacggcta catatttcgc tgcaagagcc tactatggta aatactttga 60
 ctacgaattc 70

<210> 52
 <211> 49
 <212> DNA
 <213> Artificial Sequence

<220>
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 <223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

<400> 52
 gaattcgtag tcaaagtatt taccatagta ggctcttgca gcaaatatg 49

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<210> 53
<211> 81
<212> DNA
<213> Artificial Sequence

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<221> misc_feature
<223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

<400> 53
tagcctgtgt ctcatttttt gaggttggtg atctgcaa at aggcagtgc tggcagaggtt 60
tccaaagaga aggcaaaccg t 81

<210> 54
<211> 80
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

<400> 54
cccttgaagt catcagcata tggttggtct ccagtgtagg tgtttatcca gcccatccac 60
tttaaaccct ttcctggagc 80

<210> 55
<211> 81
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

<400> 55
ctgcttcacc cagttcattc catagtttgt gaagggtatac ccagaagcct tagcggagat 60
cttgactgtc tctccaaggc t 81

<210> 56
<211> 100
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

<400> 56
tcttcagctc aggtccagac tgcaccaact ggatctgtgc ttgggcaatt tcggcagctg 60
ccatcaggaa tagcaaggtc cacaccaag ccatgaattc 100

<210> 57
<211> 63
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

<400> 57

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agtattgtga tgacccagac tcccaaattc ctgcttgtat cagcaggaga cagggttacc 60
 ata. 63

<210> 58
 <211> 64
 <212> DNA
 <213> Artificial Sequence

<220>
 <221> misc_feature
 <223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

<400> 58
 acctgcaagg ccagtcagag tgtgagtaat gatgtagctt ggtaccaaca gaaaaccagg 60
 gcag 64

<210> 59
 <211> 69
 <212> DNA
 <213> Artificial Sequence

<220>
 <221> misc_feature
 <223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 59
 tctcctaaac tgctgatata ctatgcatcc aatcgctaca ctggagtccc tgatcgcttc 60
 actggcagt 69

<210> 60
 <211> 64
 <212> DNA
 <213> Artificial Sequence

<220>
 <221> misc_feature
 <223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 60
 ggatatggga cggatttcac tttcaccatc agcaactgtgc aaggctgaag acctggcagt 60
 ttat 64

<210> 61
 <211> 69
 <212> DNA
 <213> Artificial Sequence

<220>
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 <223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 61
 ttctgycagc aggattatag ctctccgctc accttcggtg ctgggaccaa gctggacctg 60
 aaagaattc 69

<210> 62
 <211> 78
 <212> DNA
 <213> Artificial Sequence

<220>
 <221> misc_feature

<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 62
gaattctttc agctccagct tgggtcccagc accgaacgtg agcggagagc tataatcctg 60
ctgacagaaa taaactgc 78

<210> 63
<211> 63
<212> DNA
<213> Artificial Sequence

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<221> misc_feature
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<400> 63
caggtcttca gcctgcacag tgctgatggt gaaagtgaag tccgtcccat atccactgcc 60
agt 63

<210> 64
<211> 69
<212> DNA
<213> Artificial Sequence

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<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 64
gaagcgatca gggactccag tgtagcgatt ggatgcatag tatatcagca gtttaggaga 60
ctgccctgg 69

<210> 65
<211> 63
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 65
tttctgttgg taccaagcta catcattact cacactctga ctggccttgc tggttatggt 60
aac 63

<210> 66
<211> 63
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 66
cctgtctcct gctcatacaa gcaggaattt gggagtctgg gtcatacaaa tacttgcttg 60
ggc 63

<210> 67
<211> 68
<212> DNA
<213> Artificial Sequence

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<220>
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<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 67
ttcgctcagc aggattatag ctctccgctc acgttcggtg ctgggaccaa gctggagctg      60
aaagaatc                                         68

<210> 68
<211> 78
<212> DNA
<213> Artificial Sequence

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<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 68
gaattctttc agctccagct tgggtcccagc accgaacgtg agcggagagc tataatcctg      60
ctgagcgaaa taaactgc                                         78

<210> 69
<211> 399
<212> DNA
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<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<220>
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<400> 69
atg gct tgg gtg tgg acc ttg cta ttc ctg atg gca gct gcc caa agt      48
Met Ala Trp Val Trp Thr Leu Leu Phe Leu Met Ala Ala Ala Gln Ser
1 5 10 15
gcc caa gca gac att gtg atg tca cag tct cca tcc tcc cta gct gtg      96
Ala Gln Ala Asp Ile Val Met Ser Gln Ser Pro Ser Ser Leu Ala Val
20 25 30

tca gtt gga gag aag gtt act atg agc tgc aag tcc agt cag agc ctt      144
Ser Val Gly Glu Lys Val Thr Met Ser Cys Lys Ser Ser Gln Ser Leu
35 40 45
tta tat agt agc aat caa aag atc tac ttg gcc tgg tac cag cag aaa      192
Leu Tyr Ser Ser Asn Gln Lys Ile Tyr Leu Ala Trp Tyr Gln Gln Lys
50 55 60
cca ggg cag tct cct aaa ctg ctg att tac tgg gca tcc act agg gaa      240
Pro Gly Gln Ser Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu
65 70 75 80
tct ggg gtc cct gat cgc ttc aca ggc ggt gga tct ggg aca gat ttc      288
Ser Gly Val Pro Asp Arg Phe Thr Gly Gly Gly Ser Gly Thr Asp Phe
85 90 95
act ctc acc atc agc agt gtg aag gct gaa gac ctg gca gtt tat tac      336
Thr Leu Thr Ile Ser Ser Val Lys Ala Glu Asp Leu Ala Val Tyr Tyr
100 105 110
tgt cag caa tat tat aga tat cct cgg acg ttc ggt gga ggc acc aag      384
Cys-Gln-Gln-Tyr Tyr-Arg-Tyr-Pro-Arg Thr Phe Gly Gly Gly Thr Lys
115 120 125
ctg gaa atc aaa cgg                                         399

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Leu Glu Ile Lys Arg
130

<210> 70

<211> 133

<212> PRT

<213> Artificial Sequence

<220>

<221> misc_feature

<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 70

Met Ala Trp Val Trp Thr Leu Leu Phe Leu Met Ala Ala Ala Gln Ser
1 5 10 15

Ala Gln Ala Asp Ile Val Met Ser Gln Ser Pro Ser Ser Leu Ala Val
20 25 30

Ser Val Gly Glu Lys Val Thr Met Ser Cys Lys Ser Ser Gln Ser Leu
35 40 45

Leu Tyr Ser Ser Asn Gln Lys Ile Tyr Leu Ala Trp Tyr Gln Gln Lys
50 55 60

Pro Gly Gln Ser Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu
65 70 75 80

Ser Gly Val Pro Asp Arg Phe Thr Gly Gly Gly Ser Gly Thr Asp Phe
85 90 95

Thr Leu Thr Ile Ser Ser Val Lys Ala Glu Asp Leu Ala Val Tyr Tyr
100 105 110

Cys Gln Gln Tyr Tyr Arg Tyr Pro Arg Thr Phe Gly Gly Gly Thr Lys
115 120 125

Leu Glu Ile Lys Arg
130